RAW SEQUENCE LISTING PATENT APPLICATION US/09/113,924

DATE: 09/02/98 TIME: 14:23:27

INPUT SET: S28287.raw

This Raw Listing contains the General Information Section and up to the first 5 pages.

1 2		SEQUENCE LISTING ENTERE
3 4	(1)	General Information:
5 6 7		(i) APPLICANT: Brigstock, David R. Harding, Paul H.
8 9 10		(ii) TITLE OF INVENTION: HEPARIN BINDING GROWTH FACTOR (HBGF) POLYPEPTIDES
11 12		(iii) NUMBER OF SEQUENCES: 22
13		(iv) CORRESPONDENCE ADDRESS:
14		(A) ADDRESSEE: Fish & Richardson P.C.
15		(B) STREET: 4225 Executive Square, Suite 1400
16		(C) CITY: La Jolla
17		(D) STATE: CA
18		(E) COUNTRY: USA
19		(F) ZIP: 92037
20		
21		(V) COMPUTER READABLE FORM:
22		(A) MEDIUM TYPE: Diskette
23		(B) COMPUTER: IBM Compatible
24		(C) OPERATING SYSTEM: Windows 95
25		(D) SOFTWARE: FastSEQ for Windows Version 2.0b
26		
27		(vi) CURRENT APPLICATION DATA:
28		(A) APPLICATION NUMBER:
29		(B) FILING DATE:
30		
31		(vii) PRIOR APPLICATION DATA:
32		(A) APPLICATION NUMBER: 08/908,526
33		(B) FILING DATE: 07-AUG-1997
34		• •
35		
36		(viii) ATTORNEY/AGENT INFORMATION:
37		(A) NAME: Haile, Lisa A., Ph.D.
38		(B) REGISTRATION NUMBER: 38,347
39		(C) REFERENCE/DOCKET NUMBER: 08766/003001
40		(-,
41		(ix) TELECOMMUNICATION INFORMATION:
42		(A) TELEPHONE: 619/678-5070
43		(B) TELEFAX: 619/678-5099
44		\-\\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \
45		
46		(2) INFORMATION FOR SEQ ID NO:1:
		\-\

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```
47
            (i) SEQUENCE CHARACTERISTICS:
48
49
              (A) LENGTH: 16 amino acids
              (B) TYPE: amino acid
50
              (D) TOPOLOGY: linear
51
52
            (ii) MOLECULE TYPE: peptide
53
54
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
55
56
57
     Glu Glu Asn Ile Lys Lys Gly Lys Lys Xaa Ile Arg Thr Pro Lys Ile
58
                                             10
59
60
61
               (2) INFORMATION FOR SEQ ID NO:2:
62
63
            (i) SEQUENCE CHARACTERISTICS:
              (A) LENGTH: 12 amino acids
64
65
              (B) TYPE: amino acid
66
              (D) TOPOLOGY: linear
67
            (ii) MOLECULE TYPE: peptide
68
69
70
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
71
72
     Glu Asn Ile Lys Lys Gly Lys Lys Xaa Ile Arg Thr
73
74
75
76
               (2) INFORMATION FOR SEQ ID NO:3:
77
78
            (i) SEQUENCE CHARACTERISTICS:
              (A) LENGTH: 41 base pairs
79
80
              (B) TYPE: nucleic acid
81
              (C) STRANDEDNESS: single
82
              (D) TOPOLOGY: linear
83
            (ii) MOLECULE TYPE: oligonucleotide
84
85
86
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
87
     GCCGTCTAGA GCGGCCGCAT GGAAGAGAC ATTAAGAAGG G
                                                                                41
88
89
90
91
               (2) INFORMATION FOR SEQ ID NO:4:
92
93
            (i) SEQUENCE CHARACTERISTICS:
94
              (A) LENGTH: 30 base pairs
95
              (B) TYPE: nucleic acid
96
              (C) STRANDEDNESS: single
97
              (D) TOPOLOGY: linear
98
99
            (ii) MOLECULE TYPE: oligonucleotide
```

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100
101
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
102
      CCTCTGTACC GTACTTAAGC GCCGGCGACC
                                                                               30
103
104
105
                (2) INFORMATION FOR SEQ ID NO:5:
106
107
             (i) SEQUENCE CHARACTERISTICS:
               (A) LENGTH: 14 amino acids
108
109
               (B) TYPE: amino acid
               (D) TOPOLOGY: linear
110
111
112
             (ii) MOLECULE TYPE: peptide
113
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:
114
115
      Glu Glu Asn Ile Lys Lys Gly Lys Lys Cys Ile Arg Thr Pro
116
117
118
                (2) INFORMATION FOR SEQ ID NO:6:
119
120
             (i) SEQUENCE CHARACTERISTICS:
121
               (A) LENGTH: 9 amino acids
122
               (B) TYPE: amino acid
123
               (D) TOPOLOGY: linear
124
125
126
             (ii) MOLECULE TYPE: peptide
127
128
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
129
130
      Glu Glu Asn Ile Lys Lys Gly Lys Lys
131
                         5
132
133
                (2) INFORMATION FOR SEQ ID NO:7:
134
135
             (i) SEQUENCE CHARACTERISTICS:
136
               (A) LENGTH: 16 amino acids
137
               (B) TYPE: amino acid
138
               (D) TOPOLOGY: linear
139
140
             (ii) MOLECULE TYPE: peptide
141
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:
142
143
      Ile Arg Thr Pro Lys Ile Ser Lys Pro Ile Lys Phe Glu Leu Ser Gly
144
145
        1
                         5
                                             10
146
147
                (2) INFORMATION FOR SEQ ID NO:8:
148
149
150
             (i) SEQUENCE CHARACTERISTICS:
               (A) LENGTH: 17 amino acids
151
152
               (B) TYPE: amino acid
```

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```
153
               (D) TOPOLOGY: linear
154
155
             (ii) MOLECULE TYPE: peptide
156
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:
157
158
       Thr Pro Lys Ile Ser Lys Pro Ile Lys Phe Glu Leu Ser Gly Cys Thr
159
160
                         5
                                             10
       Ser
161
162
163
164
                (2) INFORMATION FOR SEQ ID NO:9:
165
166
             (i) SEQUENCE CHARACTERISTICS:
               (A) LENGTH: 10 amino acids
167
168
               (B) TYPE: amino acid
              (D) TOPOLOGY: linear
169
170
             (ii) MOLECULE TYPE: peptide
171
172
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:
173
174
       Thr Ser Met Lys Thr Tyr Arg Ala Lys Phe
175
176
        1
                         5
177
                (2) INFORMATION FOR SEQ ID NO:10:
178
179
             (i) SEQUENCE CHARACTERISTICS:
180
181
               (A) LENGTH: 13 amino acids
               (B) TYPE: amino acid
182
183
               (D) TOPOLOGY: linear
184
185
             (ii) MOLECULE TYPE: peptide
186
187
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:
188
       Thr Ser Met Lys Thr Tyr Arg Ala Lys Phe Cys Gly Val
189
190
        1
                         5
191
                (2) INFORMATION FOR SEQ ID NO:11:
192
193
194
             (i) SEQUENCE CHARACTERISTICS:
195
               (A) LENGTH: 7 amino acids
               (B) TYPE: amino acid
196
               (D) TOPOLOGY: linear
197
198
199
             (ii) MOLECULE TYPE: peptide
200
201
             (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:
202
203
       Gly Val Cys Thr Asp Gly Arg
204
205
```

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207	(2) INF	ORMATION FOR SEO ID NO:12:
208	` ,	-
209	(i) SEOUEN	CE CHARACTERISTICS:
210	, ,	TH: 8 amino acids
211	• • •	: amino acid
212	* *	LOGY: linear
213	(2) 1010	Logi. Illiear
214	(ii) MOIEC	ULE TYPE: peptide
214	(II) MOLEC	one life. peptide
	(i) CROUE	NOT DECORTOMICAL ORO ID NO.10.
216	(XI) SEQUE	NCE DESCRIPTION: SEQ ID NO:12:
217	Glas Wall Gass mb-	Arm Clas Arm Com
218	Gly Val Cys Thr	_
219	1	5
220		
221		
222	(2) INF	ORMATION FOR SEQ ID NO:13:
223	_	
224	, , –	CE CHARACTERISTICS:
225	(A) LENG	TH: 14 amino acids
226	(B) TYPE	: amino acid
227	(D) TOPO	LOGY: linear
228		•
229	(ii) MOLEC	ULE TYPE: peptide
230		
231	(xi) SEQUE	NCE DESCRIPTION: SEQ ID NO:13:
232		
233	Cys Thr Pro His	Arg Thr Thr Leu Pro Val Glu Phe Lys
234	~	
	1	5 10
235	1	5
235 236	1	5
236		ORMATION FOR SEQ ID NO:14:
236 237	(2) INF	ORMATION FOR SEQ ID NO:14:
236 237 238 239	(2) INF (i) SEQUEN	ORMATION FOR SEQ ID NO:14:
236 237 238 239 240	(2) INF (i) SEQUEN (A) LENG	ORMATION FOR SEQ ID NO:14: CE CHARACTERISTICS: TH: 13 amino acids
236 237 238 239 240 241	(2) INF (i) SEQUEN (A) LENG (B) TYPE	ORMATION FOR SEQ ID NO:14: CE CHARACTERISTICS: TH: 13 amino acids : amino acid
236 237 238 239 240 241 242	(2) INF (i) SEQUEN (A) LENG (B) TYPE	ORMATION FOR SEQ ID NO:14: CE CHARACTERISTICS: TH: 13 amino acids
236 237 238 239 240 241 242 243	(2) INF (i) SEQUEN (A) LENG (B) TYPE (D) TOPO	ORMATION FOR SEQ ID NO:14: CE CHARACTERISTICS: TH: 13 amino acids : amino acid LOGY: linear
236 237 238 239 240 241 242 243 244	(2) INF (i) SEQUEN (A) LENG (B) TYPE (D) TOPO	ORMATION FOR SEQ ID NO:14: CE CHARACTERISTICS: TH: 13 amino acids : amino acid
236 237 238 239 240 241 242 243 244 245	(2) INF (i) SEQUEN (A) LENG (B) TYPE (D) TOPO (ii) MOLEC	ORMATION FOR SEQ ID NO:14: CE CHARACTERISTICS: TH: 13 amino acids : amino acid LOGY: linear ULE TYPE: peptide
236 237 238 239 240 241 242 243 244 245 246	(2) INF (i) SEQUEN (A) LENG (B) TYPE (D) TOPO (ii) MOLEC	ORMATION FOR SEQ ID NO:14: CE CHARACTERISTICS: TH: 13 amino acids : amino acid LOGY: linear
236 237 238 239 240 241 242 243 244 245 246 247	(2) INF (i) SEQUEN (A) LENG (B) TYPE (D) TOPO (ii) MOLEC (xi) SEQUE	ORMATION FOR SEQ ID NO:14: CE CHARACTERISTICS: TH: 13 amino acids : amino acid LOGY: linear ULE TYPE: peptide NCE DESCRIPTION: SEQ ID NO:14:
236 237 238 239 240 241 242 243 244 245 246 247 248	(2) INF (i) SEQUEN (A) LENG (B) TYPE (D) TOPO (ii) MOLEC (xi) SEQUE	ORMATION FOR SEQ ID NO:14: CE CHARACTERISTICS: TH: 13 amino acids : amino acid LOGY: linear ULE TYPE: peptide NCE DESCRIPTION: SEQ ID NO:14: Thr Thr Thr Leu Pro Val Glu Phe Lys
236 237 238 239 240 241 242 243 244 245 246 247 248 249	(2) INF (i) SEQUEN (A) LENG (B) TYPE (D) TOPO (ii) MOLEC (xi) SEQUE	ORMATION FOR SEQ ID NO:14: CE CHARACTERISTICS: TH: 13 amino acids : amino acid LOGY: linear ULE TYPE: peptide NCE DESCRIPTION: SEQ ID NO:14:
236 237 238 239 240 241 242 243 244 245 246 247 248 249 250	(2) INF (i) SEQUEN (A) LENG (B) TYPE (D) TOPO (ii) MOLEC (xi) SEQUE	ORMATION FOR SEQ ID NO:14: CE CHARACTERISTICS: TH: 13 amino acids : amino acid LOGY: linear ULE TYPE: peptide NCE DESCRIPTION: SEQ ID NO:14: Thr Thr Thr Leu Pro Val Glu Phe Lys
236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251	(2) INF (i) SEQUEN (A) LENG (B) TYPE (D) TOPO (ii) MOLEC (xi) SEQUE Thr Pro His Arg	ORMATION FOR SEQ ID NO:14: CE CHARACTERISTICS: TH: 13 amino acids : amino acid LOGY: linear ULE TYPE: peptide NCE DESCRIPTION: SEQ ID NO:14: Thr Thr Thr Leu Pro Val Glu Phe Lys 5
236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252	(2) INF (i) SEQUEN (A) LENG (B) TYPE (D) TOPO (ii) MOLEC (xi) SEQUE Thr Pro His Arg	ORMATION FOR SEQ ID NO:14: CE CHARACTERISTICS: TH: 13 amino acids : amino acid LOGY: linear ULE TYPE: peptide NCE DESCRIPTION: SEQ ID NO:14: Thr Thr Thr Leu Pro Val Glu Phe Lys
236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252 253	(2) INF (i) SEQUEN (A) LENG (B) TYPE (D) TOPO (ii) MOLEC (xi) SEQUE Thr Pro His Arg 1	ORMATION FOR SEQ ID NO:14: CE CHARACTERISTICS: TH: 13 amino acids : amino acid LOGY: linear ULE TYPE: peptide NCE DESCRIPTION: SEQ ID NO:14: Thr Thr Thr Leu Pro Val Glu Phe Lys 5 10 ORMATION FOR SEQ ID NO:15:
236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252 253	(2) INF (i) SEQUEN (A) LENG (B) TYPE (D) TOPO (ii) MOLEC (xi) SEQUE Thr Pro His Arg 1 (2) INF (i) SEQUEN	ORMATION FOR SEQ ID NO:14: CE CHARACTERISTICS: TH: 13 amino acids : amino acid LOGY: linear ULE TYPE: peptide NCE DESCRIPTION: SEQ ID NO:14: Thr Thr Thr Leu Pro Val Glu Phe Lys 5 10 ORMATION FOR SEQ ID NO:15: CE CHARACTERISTICS:
236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252 253 254 255	(2) INF (i) SEQUEN (A) LENG (B) TYPE (D) TOPO (ii) MOLEC (xi) SEQUE Thr Pro His Arg 1 (2) INF (i) SEQUEN (A) LENG	ORMATION FOR SEQ ID NO:14: CE CHARACTERISTICS: TH: 13 amino acids : amino acid LOGY: linear ULE TYPE: peptide NCE DESCRIPTION: SEQ ID NO:14: Thr Thr Thr Leu Pro Val Glu Phe Lys 5 10 ORMATION FOR SEQ ID NO:15: CE CHARACTERISTICS: TH: 18 amino acids
236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252 253 254 255 256	(2) INF (i) SEQUEN (A) LENG (B) TYPE (D) TOPO (ii) MOLEC (xi) SEQUE Thr Pro His Arg 1 (2) INF (i) SEQUEN (A) LENG (B) TYPE	ORMATION FOR SEQ ID NO:14: CE CHARACTERISTICS: TH: 13 amino acids: amino acid LOGY: linear ULE TYPE: peptide NCE DESCRIPTION: SEQ ID NO:14: Thr Thr Thr Leu Pro Val Glu Phe Lys 5 10 ORMATION FOR SEQ ID NO:15: CE CHARACTERISTICS: TH: 18 amino acids : amino acid
236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252 253 254 255	(2) INF (i) SEQUEN (A) LENG (B) TYPE (D) TOPO (ii) MOLEC (xi) SEQUE Thr Pro His Arg 1 (2) INF (i) SEQUEN (A) LENG (B) TYPE	ORMATION FOR SEQ ID NO:14: CE CHARACTERISTICS: TH: 13 amino acids : amino acid LOGY: linear ULE TYPE: peptide NCE DESCRIPTION: SEQ ID NO:14: Thr Thr Thr Leu Pro Val Glu Phe Lys 5 10 ORMATION FOR SEQ ID NO:15: CE CHARACTERISTICS: TH: 18 amino acids

SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/09/113,924

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Original Text